

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/860,844DATE: 09/10/1999
TIME: 14:36:49

INPUT SET: S33270.raw

This Raw Listing contains the General
Information Section and up to the first 5 pages.

SEQUENCE LISTING

ENTERED

1
2
3 (1) General Information:
4
5 (i) APPLICANT: Weininger, Susan
6 Weininger, Arthur M
7
8 (ii) TITLE OF INVENTION: METHOD OF DETECTION OF NUCLEIC ACIDS WITH A
9 SPECIFIC SEQUENCE COMPOSITION
10
11 (iii) NUMBER OF SEQUENCES: 118
12
13 (iv) CORRESPONDENCE ADDRESS:
14 (A) ADDRESSEE: Saliwanchik & Saliwanchik
15 (B) STREET: 2421 N.W. 41st St., Suite A-1
16 (C) CITY: Gainesville
17 (D) STATE: Florida
18 (E) COUNTRY: USA
19 (F) ZIP: 32606
20
21 (v) COMPUTER READABLE FORM:
22 (A) MEDIUM TYPE: Floppy disk
23 (B) COMPUTER: IBM PC compatible
24 (C) OPERATING SYSTEM: PC-DOS/MS-DOS
25 (D) SOFTWARE: PatentIn Release #1.0, Version #1.25
26
27 (vi) CURRENT APPLICATION DATA:
28 (A) APPLICATION NUMBER: 08/860,844
29 (B) FILING DATE: 09-JUN-1997
30 (C) CLASSIFICATION:
31
32 (vii) PRIOR APPLICATION DATA:
33 (A) APPLICATION NUMBER: 08/353,476
34 (B) FILING DATE: 09-DEC-1994
35 (C) CLASSIFICATION:
36
37 (viii) ATTORNEY/AGENT INFORMATION:
38 (A) NAME: Saliwanchik, David R.
39 (B) REGISTRATION NUMBER: 35,746
40 (C) REFERENCE/DOCKET NUMBER: GP-100C1
41
42 (ix) TELECOMMUNICATION INFORMATION:
43 (A) TELEPHONE: (352) 375-8100
44 (B) TELEFAX: (352) 372-5800
45
46

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47 (2) INFORMATION FOR SEQ ID NO:1:

48

49 (i) SEQUENCE CHARACTERISTICS:

50 (A) LENGTH: 13 base pairs

51 (B) TYPE: nucleic acid

52 (C) STRANDEDNESS: both

53 (D) TOPOLOGY: linear

54

55 (ii) MOLECULE TYPE: cDNA

56

57 (iii) HYPOTHETICAL: NO

58

59 (iv) ANTI-SENSE: NO

60

61 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

62

63 TGGGGATTCC CCA

64

65

13

66 (2) INFORMATION FOR SEQ ID NO:2:

67

68 (i) SEQUENCE CHARACTERISTICS:

69 (A) LENGTH: 13 base pairs

70 (B) TYPE: nucleic acid

71 (C) STRANDEDNESS: both

72 (D) TOPOLOGY: linear

73

74 (ii) MOLECULE TYPE: cDNA

75

76 (iii) HYPOTHETICAL: NO

77

78 (iv) ANTI-SENSE: NO

79

80 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

81

82 AAGGGACTTT CCC

83

84

13

85 (2) INFORMATION FOR SEQ ID NO:3:

86

87 (i) SEQUENCE CHARACTERISTICS:

88 (A) LENGTH: 13 base pairs

89 (B) TYPE: nucleic acid

90 (C) STRANDEDNESS: both

91 (D) TOPOLOGY: linear

92

93 (ii) MOLECULE TYPE: cDNA

94

95 (iii) HYPOTHETICAL: NO

96

97 (iv) ANTI-SENSE: NO

98

99 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

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100
101 AGGGGACTTT CCG 13
102
103
104 (2) INFORMATION FOR SEQ ID NO:4:
105
106 (i) SEQUENCE CHARACTERISTICS:
107 (A) LENGTH: 15 base pairs
108 (B) TYPE: nucleic acid
109 (C) STRANDEDNESS: both
110 (D) TOPOLOGY: linear
111
112 (ii) MOLECULE TYPE: cDNA
113
114 (iii) HYPOTHETICAL: NO
115
116 (iv) ANTI-SENSE: NO
117
118 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:
119
120 GCTGGGGACT TTCCA 15
121
122
123 (2) INFORMATION FOR SEQ ID NO:5:
124
125 (i) SEQUENCE CHARACTERISTICS:
126 (A) LENGTH: 15 base pairs
127 (B) TYPE: nucleic acid
128 (C) STRANDEDNESS: both
129 (D) TOPOLOGY: linear
130
131 (ii) MOLECULE TYPE: cDNA
132
133 (iii) HYPOTHETICAL: NO
134
135 (iv) ANTI-SENSE: NO
136
137 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:
138
139 ACAAGGGACT TTCCG 15
140
141
142 (2) INFORMATION FOR SEQ ID NO:6:
143
144 (i) SEQUENCE CHARACTERISTICS:
145 (A) LENGTH: 13 base pairs
146 (B) TYPE: nucleic acid
147 (C) STRANDEDNESS: both
148 (D) TOPOLOGY: linear
149
150 (ii) MOLECULE TYPE: cDNA
151
152 (iii) HYPOTHETICAL: NO

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153
154 (iv) ANTI-SENSE: NO
155
156 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:
157
158 CCGGGTTTTTC CCC 13
159
160
161 (2) INFORMATION FOR SEQ ID NO:7:
162
163 (i) SEQUENCE CHARACTERISTICS:
164 (A) LENGTH: 27 base pairs
165 (B) TYPE: nucleic acid
166 (C) STRANDEDNESS: both
167 (D) TOPOLOGY: linear
168
169 (ii) MOLECULE TYPE: cDNA
170
171 (iii) HYPOTHETICAL: NO
172
173 (iv) ANTI-SENSE: NO
174
175 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:
176
177 AAGGGACTTT CCGCTGGGGA CTTTCCA 27
178
179
180 (2) INFORMATION FOR SEQ ID NO:8:
181
182 (i) SEQUENCE CHARACTERISTICS:
183 (A) LENGTH: 27 base pairs
184 (B) TYPE: nucleic acid
185 (C) STRANDEDNESS: both
186 (D) TOPOLOGY: linear
187
188 (ii) MOLECULE TYPE: cDNA
189
190 (iii) HYPOTHETICAL: NO
191
192 (iv) ANTI-SENSE: NO
193
194 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:
195
196 AAGGGACTTT CCGCTGGGGA CTTTCCG 27
197
198
199 (2) INFORMATION FOR SEQ ID NO:9:
200
201 (i) SEQUENCE CHARACTERISTICS:
202 (A) LENGTH: 26 base pairs
203 (B) TYPE: nucleic acid
204 (C) STRANDEDNESS: both
205 (D) TOPOLOGY: linear

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206
207 (ii) MOLECULE TYPE: cDNA
208
209 (iii) HYPOTHETICAL: NO
210
211 (iv) ANTI-SENSE: NO
212
213 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:
214
215 GCTGGGGACT TTCCAGGGAG GCGTGG 26
216
217
218 (2) INFORMATION FOR SEQ ID NO:10:
219
220 (i) SEQUENCE CHARACTERISTICS:
221 (A) LENGTH: 26 base pairs
222 (B) TYPE: nucleic acid
223 (C) STRANDEDNESS: both
224 (D) TOPOLOGY: linear
225
226 (ii) MOLECULE TYPE: cDNA
227
228 (iii) HYPOTHETICAL: NO
229
230 (iv) ANTI-SENSE: NO
231
232 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:
233
234 GCTGGGGACT TTCCAGGGGA GGTGTG 26
235
236
237 (2) INFORMATION FOR SEQ ID NO:11:
238
239 (i) SEQUENCE CHARACTERISTICS:
240 (A) LENGTH: 26 base pairs
241 (B) TYPE: nucleic acid
242 (C) STRANDEDNESS: both
243 (D) TOPOLOGY: linear
244
245 (ii) MOLECULE TYPE: cDNA
246
247 (iii) HYPOTHETICAL: NO
248
249 (iv) ANTI-SENSE: NO
250
25

PAGE: 1

SEQUENCE VERIFICATION REPORT
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Error

Original Text